

SEQUENCE LISTING

<110> Zankel, et al.

<120> MANUFACTURE OF HIGHLY PHOSPHORYLATED LYSOSOMAL ENZYMES AND USES THEREOF

<130> 30610/39385A

<150> US 60/542,586

<151> 2004-02-06

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<170> PatentIn version 3.3

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Leu Val Pro Arg Glu Leu Ser Gly Ser Ser Pro Val Leu Glu Glu Thr
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His Pro Ala His Gln Gln Gly Ala Ser Arg Pro Gly Pro Arg Asp Ala
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Pro Pro Asn Ser Arg Phe Asp Cys Ala Pro Asp Lys Ala Ile Thr Gln
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Glu Gln Cys Glu Ala Arg Gly Cys Cys Tyr Ile Pro Ala Lys Gln Gly
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Tyr Pro Ser Tyr Lys Leu Glu Asn Leu Ser Ser Glu Met Gly Tyr
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Phe Thr Ile Lys Asp Pro Ala Asn Arg Arg Tyr Glu Val Pro Leu Glu
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Phe Ser Glu Glu Pro Phe Gly Val Ile Val His Arg Gln Leu Asp Gly
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Arg Val Leu Leu Asn Thr Thr Val Ala Pro Leu Phe Phe Ala Asp Gln
225 230 235 240

Phe Leu Gln Leu Ser Thr Ser Leu Pro Ser Gln Tyr Ile Thr Gly Leu
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Ala Glu His Leu Ser Pro Leu Met Leu Ser Thr Ser Trp Thr Arg Ile
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Thr Leu Trp Asn Arg Asp Leu Ala Pro Thr Pro Gly Ala Asn Leu Tyr
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Gly Ser His Pro Phe Tyr Leu Ala Leu Glu Asp Gly Gly Ser Ala His
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Ser Pro Ala Leu Ser Trp Arg Ser Thr Gly Gly Ile Leu Asp Val Tyr
325 330 335

Ile Phe Leu Gly Pro Glu Pro Lys Ser Val Val Gln Gln Tyr Leu Asp
340 345 350

Val Val Gly Tyr Pro Phe Met Pro Pro Tyr Trp Gly Leu Gly Phe His
355 360 365

Leu Cys Arg Trp Gly Tyr Ser Ser Thr Ala Ile Thr Arg Gln Val Val
370 375 380

Glu Asn Met Thr Arg Ala His Phe Pro Leu Asp Val Gln Trp Asn Asp
385 390 395 400

Leu Asp Tyr Met Asp Ser Arg Arg Asp Phe Thr Phe Asn Lys Asp Gly
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Phe Arg Asp Phe Pro Ala Met Val Gln Glu Leu His Gln Gly Gly Arg
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Arg Tyr Met Met Ile Val Asp Pro Ala Ile Ser Ser Ser Gly Pro Ala
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Gly Ser Tyr Arg Pro Tyr Asp Glu Gly Leu Arg Arg Gly Val Phe Ile
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Cys Pro Asn Asn Glu Leu Glu Asn Pro Pro Tyr Val Pro Gly Val Val
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Gly Gly Thr Leu Gln Ala Ala Thr Ile Cys Ala Ser Ser His Gln Phe
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Leu Ser Thr His Tyr Asn Leu His Asn Leu Tyr Gly Leu Thr Glu Pro
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Ile Ala Ser His Arg Ala Leu Val Lys Ala Arg Gly Thr Arg Pro Phe
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Asp Val Cys Gly Phe Leu Gly Asn Thr Ser Glu Glu Leu Cys Val Arg
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Gln Ala Met Arg Lys Ala Leu Thr Leu Arg Tyr Ala Leu Leu Pro His
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Leu Tyr Thr Leu Phe His Gln Ala His Val Ala Gly Glu Thr Val Ala
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Arg Pro Leu Phe Leu Glu Phe Pro Lys Asp Ser Ser Thr Trp Thr Val
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Asp His Gln Leu Leu Trp Gly Glu Ala Leu Leu Ile Thr Pro Val Leu
740 745 750

Gln Ala Gly Lys Ala Glu Val Thr Gly Tyr Phe Pro Leu Gly Thr Trp
755 760 765

Tyr Asp Leu Gln Thr Val Pro Ile Glu Ala Leu Gly Ser Leu Pro Pro
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Pro Pro Ala Ala Pro Arg Glu Pro Ala Ile His Ser Glu Gly Gln Trp
785 790 795 800

Val Thr Leu Pro Ala Pro Leu Asp Thr Ile Asn Val His Leu Arg Ala
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Gly Tyr Ile Ile Pro Leu Gln Gly Pro Gly Leu Thr Thr Glu Ser
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